

LOCUS NP_060628 796 aa PRI 04-JUL-2000
 DEFINITION hypothetical protein FLJ10624 [Homo sapiens].
 ACCESSION NP_060628
 PID g8922557
 VERSION NP_060628.1 GI:8922557
 DBSOURCE REFSEQ: accession NM_018158.1
 KEYWORDS .
 SOURCE human.
 ORGANISM Homo sapiens
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
 REFERENCE 1 (sites)
 AUTHORS Isogai,T., Ota,T., Hayashi,K., Sugiyama,T., Otsuki,T., Suzuki,Y.,
 Nishikawa,T., Nagai,K., Sugano,S., Ishibashi,T., Fujimori,K.,
 Tanai,H., Kimata,M., Watanabe,M., Hiraoka,S., Ishii,S., Kawai,Y.,
 Saito,K., Yamamoto,J., Wakamatsu,A., Nakamura,Y., Nagahari,K.,
 Masuho,Y. and Kanehori,K.
 TITLE NEDO human cDNA sequencing project
 JOURNAL Unpublished
 COMMENT REFSEQ: The reference sequence was derived from AK001486.1.
 PREDICTED RefSeq: This reference sequence record is derived from a
 cDNA of unknown function. The existence of a cDNA provides
 experimental evidence for the gene; however, the location of the
 open reading frame and corresponding protein sequence have been
 predicted and may change as additional sequence and experimental
 data become available.
 FEATURES Location/Qualifiers
 source 1..796
 /organism="Homo sapiens"
 /db_xref="taxon:9606"
 /clone="NT2RP2005525"
 /cell_line="NT2"
 /cell_type="teratocarcinoma"
 /clone_lib="NT2RP2"
 /note="cloning vector: pME18SFL3; mRNA from NT2 neuronal
 precursor cells after 2-weeks retinoic acid (RA)
 induction."
 Protein 1..796
 /product="hypothetical protein FLJ10624"
 CDS 1..796
 /gene="FLJ10624"
 /coded_by="NM_018158.1:283..2673"
 ORIGIN
 1 mlaplrnapg regatspspp tdatgslgew dvdrnvkteq wvskeriskl hrlrmadils
 61 qsetlasqdl sgdfkkpalp vspaarskap assssnpeev qkegptalqd snsgepdipp
 121 pqpdcdgdfrs lqeeqsrptt avsspggpar appyqeppwg gpataapysle tlkggttilgt
 181 rslkgtsycl fgrlsgcdvc lehpsvsryh avlqhrasgp dgecdsngpg fylydulgsth
 241 gtflnktrip prtycrvhvg hvvrfggstr lfilqgpeed reaeseltvt qlkelrkqqq
 301 illekkmrlre dsdeeeeemd serkinagsq ddemgctwgm gedaveddae enpivlefqq
 361 ereafyikdp kkvlgqffdr egeeleyefd eqghstwlcr vrlpvddstg kqlvaeaihs
 421 gkkkeamiqc sleacrilidt lglirqeavk rkrkaknwed edfydsddd flidrtglik
 481 krlnrmkkag kidekpetfe slvaklnae relseiserl kassqvlses psqdsldafm
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 601 kkltlplfga mkggskfk1k tgtvgklppk rpelpptlmr mkdepeveee eeeeeeeeke
 661 keehekkkle dgslsrpqpe iepeaaavqem rpptdlthfk etqthenmsq lseeeeqnky
 721 qdcskttslc agpsaskney eksrgelkk ktpgpgklpp tlsskypedd pdycvwvppe
 781 gqsgdgrthl ndkygy

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